



cket No.: PF-0539 USN

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of:

Lal et al.

Title:

PROTEINS REGULATING GENE EXPRESSION

Serial No.:

09/701,674

Filing Date:

To Be Assigned

Examiner:

To Be Assigned

Group Art Unit:

To Be Assigned

United States Patent and Trademark Office

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SUBSTITUTE SUBMISSION UNDER 37 CFR §1.821-1.825 SEQUENCE LISTING

Sir:

In accordance with the requirements of 37 CFR \$1.821-1.825, Applicants hereby submit one (1) substitute diskette containing the computer-readable information for the "Sequence Listing" of the above-identified application. The substitute diskette complies with the requirements of 37 CFR §1.824 and is IBM PC compatible using a Windows NT Operating System with WordPerfect software and saved in ASCII text format.

Enclosed is a paper copy of the Substitute Sequence Listing.

The content of the Substitute Sequence Listing paper copy is identical to the computer-readable copy, as required under 37 CFR § 1.821(f). No new matter has been added.

Respectfully submitted.

INCYTE GENOMICS, INC.

Date: Folial 2002

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102876

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09/701,674



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/701,674
Source:	, PUTOGRUSH
Date Processed by STIC:	8/26/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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Revised 01/29/2002



PN/09

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PATENT APPLICATION: US/09/701,674

DATE: 08/26/2002

TIME: 13:40:35

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            LAL, Preeti
            YUE, Henry
            TANG, Y. Tom
    7
            HILLMAN, Jennifer L.
    Я
            BANDMAN, Olga
    9
   10
            CORLEY, Neil C.
                                                                    Does Not Comply
            GUEGLER, Karl J.
   11
                                                                Corrected Diskette Needec
   12
            GORGONE, Gina A.
            BAUGHN, Mariah R.
   13
   14
            PATTERSON, Chandra
           LU, Dyung Aina M.
   15
   17 <120> TITLE OF INVENTION: PROTEINS REGULATING GENE EXPRESSION
   19 <130> FILE REFERENCE: PF-0539 PCT
-> 21 <140> CURRENT APPLICATION NUMBER: US/09/701,674
  22 <141> CURRENT FILING DATE: 2002-08-26
   24 <150> PRIOR APPLICATION NUMBER: 60/089,029; 60/094,575;)60/104,624
                                              place prior bist application runter separately directly above its
-> 25 <151> PRIOR FILING DATE: (1998-06-12+ 1998-07-29;) 1998-10-14
   27 <160> NUMBER OF SEQ ID NOS: 62
   29 <170> SOFTWARE: PERL Program
RORED SEQUENCES
   2400 <210> SEQ ID NO: 36
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   2403 <213> ORGANISM: Homo sapiens
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DATE: 08/26/2002

PATENT APPLICATION: US/09/701,674

TIME: 13:40:35

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-> 2451 2418

Use of providing Xna has been detected in the Sequence Listing.

Review 1. Acquence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of

each sequence using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/701,674
DATE: 08/26/2002
TIME: 13:40:36

Input Set : A:\pf-0539usn_seqlist.txt
Output Set: N:\CRF3\08262002\I701674.raw

21 M:270 C: Current Application Number differs, Replaced Current Application Number 22 M:271 C: Current Filing Date differs, Replaced Current Filing Date 25 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD 281 M:283 W: Missing Blank Line separator, <220> field identifier 319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:210 2356 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2358 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2360 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2362 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2364 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2366 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2374 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:35 2375 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0 2376 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:60 2389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:840 2390 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:900 2391 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:960 2392 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1020 2393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1080 2394 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1140 2395 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1200 2396 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1260 2397 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1320 2450 M:254 E. No. of Bases conflict, LENGTH: Input: 0 Counted: 2418 SEQ:36 2451 M:252 E: No. of Seq. differs, <211> LENGTH:Input:2402 Found:2418 SEQ:36 2480 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:660 2481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:720 2482 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:780 2483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:840 2697 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:2220 2813 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:60 2814 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:120 2815 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:180 3153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:1920

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